46

(2) INFORMATION FOR SEQ ID NO:1:

RAW SEQUENCE LISTING PATENT APPLICATION US/09/033,662

DATE: 08/08/98 TIME: 15:52:36

INPUT SET: S27966.raw

This Raw Listing contains the General Information Section and up to the first 5 pages.

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ENTERED
 1
                                       SEQUENCE LISTING
 2
           General Information:
 3
    (1)
 4
          (i) APPLICANT: HU, JING-SHAN
 5
 6
                         OLSEN, HENRIK S.
 7
                         ROSEN, CRAIG A.
 8
        (ii) TITLE OF INVENTION: HUMAN VASCULAR ENDOTHELIAL GROWTH FACTOR 3
 9
10
        (iii) NUMBER OF SEQUENCES: 8
11
12
         (iv) CORRESPONDENCE ADDRESS:
13
               (A) ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX P.L.L.C.
14
15
               (B) STREET: 1100 NEW YORK AVE, NW
16
               (C) CITY: WASHINGTON
               (D) STATE: DC
17
18
               (E) COUNTRY: USA
19
               (F) ZIP: 20005
20
          (V) COMPUTER READABLE FORM:
21
               (A) MEDIUM TYPE: Floppy disk
22
               (B) COMPUTER: IBM PC compatible
23
24
               (C) OPERATING SYSTEM: PC-DOS/MS-DOS
               (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
25
26
27
        (vi) CURRENT APPLICATION DATA:
               (A) APPLICATION NUMBER: US 09/033,662
28
29
               (B) FILING DATE: 03-MAR-1998
30
               (C) CLASSIFICATION:
31
32
       (vii) PRIOR APPLICATION DATA:
33
               (A) APPLICATION NUMBER: US 08/469,641
34
               (B) FILING DATE: 06-JUN-1995
35
36
       (viii) ATTORNEY/AGENT INFORMATION:
37
               (A) NAME: STEFFE, ERIC K.
38
               (B) REGISTRATION NUMBER: 36,688
39
               (C) REFERENCE/DOCKET NUMBER: 1488.1040001
40
        (ix) TELECOMMUNICATION INFORMATION:
41
42
               (A) TELEPHONE: 202-371-2600
43
               (B) TELEFAX: 202-371-2540
44
45
```

E-q

RAW SEQUENCE LISTING PATENT APPLICATION US/09/033,662

DATE: 08/08/98 TIME: 15:52:37

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47																	
48	(i) SEQUENCE CHARACTERISTICS:																
49			(1	A) LI	engti	H: 66	56 ba	ase j	pair	3							
50			(1	B) T	YPE:	nuc	leic	acio	£								
51			((C) S	PRANI	DEDNI	ESS:	both	n n								
52			(1	D) T(OPOL	OGY:	botl	n n									
53																	
54		(ii)	MOI	LECUI	LE T	YPE:	CDN	A.									
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57		(ix)) FE	ATURI	E :												
58		` '	· a	A) N	AME/I	KEY:	CDS										
59			•	B) L				563									
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61																	
62		(xi)	SEC	OUEN	E DI	SCR	CPTIC	ON: S	SEO :	ED NO	0:1:						
63		(,,							-							
64	ATG	AGA	AGG	TGT	AGA	АТА	AGT	GGG	AGG	ccc	CCG	GCG	ccc	CCC	GGT	GTC	48
65															Gly		
66	1	3	5	- 2 -	5			2	5	10					15		
67	_				_												
68	CCC	GCC	CAG	GCC	ССТ	GTC	TCC	CAG	ССТ	САТ	GCC	ССТ	GGC	CAC	CAG	AGG	96
69															Gln		, ,
70				20					25	E			,	30		9	
71																	
72	ΔΔΔ	стс	GTG	ТСΔ	TGG	ΔΤΔ	САТ	GTG	тΔт	ΔСТ	CGC	CCT	ACC	TGC	CAG	CCC	144
73															Gln		111
74	Lys	· · · ·	35	DCI			пор	40	* y *		~-9	n.u	45	Cys	0111	110	
75													10				
76	CGG	GAG	стс	GTG	GTG	CCC	ጥጥር	ΔСТ	GTG	GAG	מיזימ	ΔТС	ממר	ACC	GTG	GCC	192
77															Val		-,-
78	9	50					55					60	,				
79												•					
80	ΔΔΔ	CAG	СТС	стс	ccc	AGC	TGC	GTG	ΔСТ	GTG	CAG	CGC	тат	GGT	GGC	TGC	240
81															Gly		210
82	65	01	200	·ul		70	4			,	75	9	0,5	0_1	0_3	80	
83	• • •					, 0					. •					•	
84	TCC	ССТ	GAC	СΔТ	GGC	ста	GAG	ጥርጥ	GTG	CCC	ΔСТ	GGG	CAG	CAC	CAA	GTC	288
85															Gln		200
86	Cys	110	rop	rop	85	DC G	014	Cyb	*41	90			01	****	95	V U 1	
87					05					70					,,		
88	CGG	λΤС	CAG	ልጥሮ	CTC	λπα	ልጥሮ	CGG	ጥልሮ	CCG	ACC	λСΤ	CAG	СТС	GGG	GAG	336
89															Gly		330
90	AL 9	мес	GIN	100	Беа	Mec	116	ALG	105	110	561	Der	GIII	110	Gry	GIU	
91				100					103					110			
92	y m.c.	TOO	ama	C A A	C 3 3	CAC	300	CAC	mam.	CAA	TICC.	202	COM	***	AAA	220	384
93															Lys		304
94	M C C	PET	115	GIU	GIU	1112	SEI	120	Cys	GIU	Cys	ALY.	125	пåэ	пåэ	ny 5	
95			113					120					123			•	
96	GVG	አረጣ	CCT	QTQ.	220	CCA	GNG	אממ	CCT	CCT	አ ሮሞ	ccc	CAC	CAC	CGT	ccc	432
97															Arg		432
98	voh	130	та	AGT	пya	FIO	135	AT 9	VT a	via	TILL	140	1112	117.2	wrd	FIU	
99		130					133					140					
73																	

152

RAW SEQUENCE LISTING PATENT APPLICATION US/09/033,662

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														IN	PUT	SET:	S27966. raw	,
100	CAG	ccc	CGT	TCT	GTT	CCG	GGC	TGG	GAC	TCT	GCC	CCC	GGA				4.8	
101				Ser														
102	145		_			150	-	-	_		155		-			160		
103																		
104	CCA	GCT	GAC	ATC	ACC	CAA	TCC	CAC	TCC	AGC	CCC	AGG	CCC	CTC	TGC	CCA	52	8:
105	Pro	Ala	Asp	Ile	Thr	Gln	Ser	His	Ser	Ser	Pro	Arg	Pro	Leu	Cys	Pro		
106			_		165					170					175			
107																		
108				CAG													57	6.
109	Arg	Cys	Thr	Gln	His	His	Gln	Cys	Pro	Asp	Pro	Arg	Thr	Cys	Arg	Cys		
110				180					185					190				
111																		
112				CGC													62	4
113	Arg	Cys	Arg	Arg	Arg	Ser	Phe	Leu	Arg	Cys	Gln	Gly	Arg	Gly	Leu	Glu		
114			195					200					205					
115																		
116	_			GAC										TGA			66	6
117	Leu	Asn	Pro	Asp	Thr	Cys	_	Cys	Arg	Lys	Leu	_	Arg					
118		210					215					220						
119										•								
120																		
121	(2)	INF	ORMA'	rion	FOR	SEQ	ID I	NO: 2	:									
122						a		3D T 01	aa									
123			(1):	SEQUI							_							
124				(A) LEI	NGTH	: 22.	L am:	ruo a	acias	3							
				/ D	m 1771	n a .												
125				•		PE: 8	_											
126				•		PE: 8	_											
126 127		1 :	i i \)	(D	TO	POLO	3Y:]	line	ar									
126 127 128		(:	ii) ł	•	TO	POLO	3Y:]	line	ar									
126 127 128 129		·	·	(D)) TO	POLO TYPI	3Y:] E: p:	line: rote:	ar in	O ID	NO:	2:						
126 127 128 129 130		·	·	(D) TO	POLO TYPI	3Y:] E: p:	line: rote:	ar in) ID	NO:2	2:						
126 127 128 129 130 131	Met	(2	xi) :	(D) MOLEC SEQUI) TOI CULE ENCE	POLOG TYPI DESG	BY:] E: pi	Linea rote: rion	ar in : SE(Pro	Pro	Glv	Val		
126 127 128 129 130 131		(:	xi) :	(D)) TOI CULE ENCE	POLOG TYPI DESG	BY:] E: pi	Linea rote: rion	ar in : SE(Pro	Pro	Gly 15	Val		
126 127 128 129 130 131	Met 1	(:	xi) :	(D) MOLEC SEQUI	TOICULE ENCE Arg	POLOG TYPI DESG	BY:] E: pi	Linea rote: rion	ar in : SE(Pro			Pro	Pro		Val		
126 127 128 129 130 131 132	1	(:	xi) :	(D) MOLEC SEQUI	OULE ENCE Arg	TYPI DESC	E: pr CRIPT	linea rote rion Gly	ar in : SE(Pro 10	Pro	Ala			15			
126 127 128 129 130 131 132 133	1	(:	xi) :	(D) MOLEC SEQUI	OULE ENCE Arg	TYPI DESC	E: pr CRIPT	linea rote rion Gly	ar in : SE(Pro 10	Pro	Ala			15			
126 127 128 129 130 131 132 133 134	1	(:	xi) :	(D) MOLEC SEQUI Cys	OULE ENCE Arg	TYPI DESC	E: pr CRIPT	linea rote rion Gly	ar in : SEG Arg	Pro 10	Pro	Ala		His	15			
126 127 128 129 130 131 132 133 134 135	l Pro	Arg	xi) : Arg Gln	(D) MOLEC SEQUI Cys	OULE ENCE Arg 5	TYPH DESC Ile Val	GY:] E: pr CRIPT Ser Ser	linea rote rion Gly Gln	in : SEG Arg Pro 25	Pro 10 Asp	Pro Ala	Ala	Gly	His 30	15 Gln	Arg		
126 127 128 129 130 131 132 133 134 135 136	l Pro	Arg	xi) : Arg Gln	MOLEG SEQUI Cys Ala 20	OULE ENCE Arg 5	TYPH DESC Ile Val	GY:] E: pr CRIPT Ser Ser	linea rote rion Gly Gln	in : SEG Arg Pro 25	Pro 10 Asp	Pro Ala	Ala	Gly	His 30	15 Gln	Arg		
126 127 128 129 130 131 132 133 134 135 136 137	1 Pro Lys	Arg Ala Val	Arg Gln Val	(D) MOLEG SEQUI Cys Ala 20 Ser	OULE ENCE Arg Fro Trp	TYPI DESC Ile Val	GY: G E: pr CRIP' Ser Ser	rote: FION Gly Gln Val	in : SEG Arg Pro 25	Pro 10 Asp	Pro Ala Arg	Ala Pro Ala	Gly Thr 45	His 30 Cys	15 Gln Gln	Arg Pro		
126 127 128 129 130 131 132 133 134 135 136 137 138 139 140 141	1 Pro Lys	Arg Ala Val	Arg Gln Val	MOLEG SEQUI Cys Ala 20	OULE ENCE Arg Fro Trp	TYPI DESC Ile Val	GY: G E: pr CRIP' Ser Ser	rote: FION Gly Gln Val	in : SEG Arg Pro 25	Pro 10 Asp	Pro Ala Arg	Ala Pro Ala	Gly Thr 45	His 30 Cys	15 Gln Gln	Arg Pro		
126 127 128 129 130 131 132 133 134 135 136 137 138 139 140 141 142	1 Pro Lys	Arg Ala Val	Arg Gln Val	(D) MOLEG SEQUI Cys Ala 20 Ser	OULE ENCE Arg Fro Trp	TYPI DESC Ile Val	E: pr CRIP' Ser Ser	rote: FION Gly Gln Val	in : SEG Arg Pro 25	Pro 10 Asp	Pro Ala Arg	Ala Pro Ala	Gly Thr 45	His 30 Cys	15 Gln Gln	Arg Pro		
126 127 128 129 130 131 132 133 134 135 136 137 138 139 140 141 142 143	Pro Lys	Arg Ala Val Glu 50	Arg Gln Val 35	MOLEG SEQUI Cys Ala 20 Ser	CULE ENCE Arg 5 Pro Trp	TYPI DESC Ile Val Ile	E: processor pro	rote: FION Gly Gln Val 40	in : SEG Arg Pro 25 Tyr	Pro 10 Asp Thr	Pro Ala Arg Leu	Ala Pro Ala Met 60	Gly Thr 45	His 30 Cys Thr	15 Gln Gln Val	Arg Pro		
126 127 128 129 130 131 132 133 134 135 136 137 138 139 140 141 142 143 144	Pro Lys Arg	Arg Ala Val Glu 50	Arg Gln Val 35	(D) MOLEG SEQUI Cys Ala 20 Ser	CULE ENCE Arg 5 Pro Trp	TYPI DESC Ile Val Ile Pro	E: processor pro	rote: FION Gly Gln Val 40	in : SEG Arg Pro 25 Tyr	Pro 10 Asp Thr	Pro Ala Arg Leu Gln	Ala Pro Ala Met 60	Gly Thr 45	His 30 Cys Thr	15 Gln Gln Val	Arg Pro Ala Cys		
126 127 128 129 130 131 132 133 134 135 136 137 138 139 140 141 142 143 144	Pro Lys	Arg Ala Val Glu 50	Arg Gln Val 35	MOLEG SEQUI Cys Ala 20 Ser	CULE ENCE Arg 5 Pro Trp	TYPI DESC Ile Val Ile	E: processor pro	rote: FION Gly Gln Val 40	in : SEG Arg Pro 25 Tyr	Pro 10 Asp Thr	Pro Ala Arg Leu	Ala Pro Ala Met	Gly Thr 45	His 30 Cys Thr	15 Gln Gln Val	Arg Pro		
126 127 128 129 130 131 132 133 134 135 136 137 138 139 140 141 142 143 144	Pro Lys Arg Lys 65	Arg Ala Val Glu 50 Gln	Arg Gln Val 35 Val	MOLEG SEQUI Cys Ala 20 Ser Val	CULE ENCE Arg Fro Trp Val	TYPI DESC Ile Val Ile Pro Ser 70	E: processor pro	rote: rION Gly Gln Val 40 Thr	in : SEG Arg Pro 25 Tyr Val	Pro 10 Asp Thr Glu Val	Pro Ala Arg Leu Gln 75	Ala Pro Ala Met 60 Arg	Gly Thr 45 Gly Cys	His 30 Cys Thr	Gln Cln Val	Arg Pro Ala Cys 80		
126 127 128 129 130 131 132 133 134 135 136 137 138 139 140 141 142 143 144 145 146 147	Pro Lys Arg Lys 65	Arg Ala Val Glu 50 Gln	Arg Gln Val 35 Val	MOLEG SEQUI Cys Ala 20 Ser	CULE ENCE Arg Fro Trp Val Pro Gly	TYPI DESC Ile Val Ile Pro Ser 70	E: processor pro	rote: rION Gly Gln Val 40 Thr	in : SEG Arg Pro 25 Tyr Val	Pro 10 Asp Thr Glu Val	Pro Ala Arg Leu Gln 75	Ala Pro Ala Met 60 Arg	Gly Thr 45 Gly Cys	His 30 Cys Thr	Gln Cln Val Cly Cln	Arg Pro Ala Cys 80		
126 127 128 129 130 131 132 133 134 135 136 137 138 139 140 141 142 143 144 145 146 147 148	Pro Lys Arg Lys 65	Arg Ala Val Glu 50 Gln	Arg Gln Val 35 Val	MOLEG SEQUI Cys Ala 20 Ser Val	CULE ENCE Arg Fro Trp Val	TYPI DESC Ile Val Ile Pro Ser 70	E: processor pro	rote: rION Gly Gln Val 40 Thr	in : SEG Arg Pro 25 Tyr Val	Pro 10 Asp Thr Glu Val	Pro Ala Arg Leu Gln 75	Ala Pro Ala Met 60 Arg	Gly Thr 45 Gly Cys	His 30 Cys Thr	Gln Cln Val	Arg Pro Ala Cys 80		
126 127 128 129 130 131 132 133 134 135 136 137 138 139 140 141 142 143 144 145 146 147 148	Pro Lys Arg Lys 65 Cys	Arg Ala Val Glu 50 Gln Pro	Arg Gln Val 35 Val Leu Asp	MOLEG SEQUI Cys Ala 20 Ser Val Val	CULE ENCE Arg Fro Trp Val Pro Gly 85	TYPI DESC Ile Val Ile Pro Ser 70 Leu	E: process of the control of the con	rote: rION Gly Gln Val 40 Thr Val Cys	in : SEG Arg Pro 25 Tyr Val Thr	Pro 10 Asp Thr Glu Val	Pro Ala Arg Leu Gln 75	Ala Pro Ala Met 60 Arg	Gly Thr 45 Gly Cys	His 30 Cys Thr Gly	Gln Cln Cly Cly Cln 95	Arg Pro Ala Cys 80 Val		
126 127 128 129 130 131 132 133 134 135 136 137 138 139 140 141 142 143 144 145 146 147 148	Pro Lys Arg Lys 65 Cys	Arg Ala Val Glu 50 Gln Pro	Arg Gln Val 35 Val Leu Asp	MOLEG SEQUI Cys Ala 20 Ser Val	CULE ENCE Arg Fro Trp Val Pro Gly 85	TYPI DESC Ile Val Ile Pro Ser 70 Leu	E: process of the control of the con	rote: rION Gly Gln Val 40 Thr Val Cys	in : SEG Arg Pro 25 Tyr Val Thr	Pro 10 Asp Thr Glu Val	Pro Ala Arg Leu Gln 75	Ala Pro Ala Met 60 Arg	Gly Thr 45 Gly Cys	His 30 Cys Thr Gly	Gln Cln Cly Cly Cln 95	Arg Pro Ala Cys 80 Val		

RAW SEQUENCE LISTING PATENT APPLICATION US/09/033,662

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Met Ser Leu Glu Glu His Ser Gln Cys Glu Cys Arg Pro Lys Lys
154
                                  120
155
     Asp Ser Ala Val Lys Pro Asp Arg Ala Ala Thr Pro His His Arg Pro
156
157
158
     Gln Pro Arg Ser Val Pro Gly Trp Asp Ser Ala Pro Gly Ala Pro Ser
159
                          150
160
161
     Pro Ala Asp Ile Thr Gln Ser His Ser Ser Pro Arg Pro Leu Cys Pro
162
                                           170
163
                      165
164
     Arg Cys Thr Gln His His Gln Cys Pro Asp Pro Arg Thr Cys Arg Cys
165
166
                                       185
167
     Arg Cys Arg Arg Arg Ser Phe Leu Arg Cys Gln Gly Arg Gly Leu Glu
168
169
              195
170
     Leu Asn Pro Asp Thr Cys Arg Cys Arg Lys Leu Arg Arg
171
          210
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172
173
174
175
     (2) INFORMATION FOR SEQ ID NO:3:
176
177
           (i) SEQUENCE CHARACTERISTICS:
                (A) LENGTH: 29 base pairs
178
                (B) TYPE: nucleic acid
179
180
                (C) STRANDEDNESS: both
181
                (D) TOPOLOGY: linear
182
         (ii) MOLECULE TYPE: cDNA
183
184
185
186
187
188
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:
189
190
     GCATGGATCC CAGCCTGATG CCCCTGGCC
                                                                               29
191
     (2) INFORMATION FOR SEQ ID NO:4:
192
193
194
           (i) SEQUENCE CHARACTERISTICS:
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                (A) LENGTH: 30 base pairs
196
                (B) TYPE: nucleic acid
197
                (C) STRANDEDNESS: both
198
                (D) TOPOLOGY: linear
199
          (ii) MOLECULE TYPE: cDNA
200
201
202
203
204
205
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:
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RAW SEQUENCE LISTING PATENT APPLICATION US/09/033,662

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	INPE	/1 SE1: 32/900.raw
206	CONTROL ON COCHOCHOLO MOTONING	20
207	GCATTCTAGA CCCTGCTGAG TCTGAAAAGC	30
208 209	(2) THEODMARTON FOR CEO TO NO. 5.	
210	(2) INFORMATION FOR SEQ ID NO:5:	
211	(i) SEQUENCE CHARACTERISTICS:	
212	(A) LENGTH: 29 base pairs	
213	(B) TYPE: nucleic acid	
214	(C) STRANDEDNESS: both	
215	(D) TOPOLOGY: linear	
216	(b) 1010B001. 11.10d2	
217	(ii) MOLECULE TYPE: cDNA	
218	(11) 110220022 11121 051111	
219		
220		
221		
222	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:	
223	(,,	
224	GACTGCATGC ACCAGAGGAA AGTGGTGTC	29
225		
226	(2) INFORMATION FOR SEQ ID NO:6:	
227		
228	(i) SEQUENCE CHARACTERISTICS:	
229	(A) LENGTH: 29 base pairs	
230	(B) TYPE: nucleic acid	
231	(C) STRANDEDNESS: both	
232	(D) TOPOLOGY: linear	
233		
234	(ii) MOLECULE TYPE: cDNA	
235		
236		
237		
238		
239	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:	
240		
241	GACTAGATCT CCTTCGCAGC TTCCGGCAC	29
242	•	
243	(2) INFORMATION FOR SEQ ID NO:7:	
244		
245	(i) SEQUENCE CHARACTERISTICS:	
246	(A) LENGTH: 14 amino acids	
247	(B) TYPE: amino acid	
248	(C) STRANDEDNESS: single	
249	(D) TOPOLOGY: Not Relevant	
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251	(ii) MOLECULE TYPE: peptide	
252		
253		
254		
255		
256	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:	
257		
258	Pro Xaa Cys Val Xaa Xaa Xaa Arg Cys Xaa Gly Cys Cys A	.sn

SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/09/033,662

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